

SEQUENCE LISTING

<110> Wang, Huaming

<120> Novel Phenol Oxidizing Enzymes

<130> GC567

<140> 09/218, 702

<141> 1998-12-22

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<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1791

<212> DNA

<213> Stachybotrys sp.

<400> 1

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<210> 2

<211> 594

<212> PRT

<213> *Stachybotrys* sp.

<400> 2

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35 40 45
Ala Ala Gly Asp Asp Asp Trp Glu Ser Pro Pro Tyr Asn Leu Leu Tyr
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Arg Asn Ala Leu Pro Ile Pro Pro Val Lys Gln Pro Lys Met Ile Ile
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Thr Asn Pro Val Thr Gly Lys Asp Ile Trp Tyr Tyr Glu Ile Glu Ile
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Val Gly Tyr Asp Gly Met Ser Pro Gly Pro Thr Phe Asn Val Pro Arg
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Asn Asp Glu Ala Glu Asp Ala Leu Gly Leu Pro Ser Gly Tyr Gly Glu
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Arg Phe Val Val Ser Ser Gly Thr Val Glu Asp Asn Ser Gln Val Pro
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Ser Thr Leu Arg Asp Val Pro Phe Pro His Lys Glu Gly Pro Ala
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Leu Gly Thr Val Glu Val Trp Glu Leu Glu Asn Ser Ser Gly Gly Trp
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<210> 3
 <211> 3677
 <212> DNA
 <213> *Stachybotrys chartarum*

<400> 3

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<211> 568

<212> PRT

<213> Bilirubin oxidase

<400> 4

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Gly	His	Leu	Phe	Lys	Arg	Val	Ala	Gln	Ile	Ser	Pro	Gln	Tyr	Pro	Met
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Phe	Thr	Val	Pro	Leu	Pro	Ile	Pro	Pro	Val	Lys	Gln	Pro	Arg	Leu	Thr
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Val	Thr	Asn	Pro	Val	Asn	Gly	Gln	Glu	Ile	Trp	Tyr	Tyr	Glu	Val	Glu
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Ile	Lys	Pro	Phe	Thr	His	Gln	Val	Tyr	Pro	Asp	Leu	Gly	Ser	Ala	Asp
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Leu	Val	Gly	Tyr	Asp	Gly	Met	Ser	Pro	Gly	Pro	Thr	Phe	Gln	Val	Pro
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Arg	Gly	Val	Glu	Thr	Val	Val	Arg	Phe	Ile	Asn	Asn	Ala	Glu	Ala	Pro
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Asn Ser Val His Leu His Gly Ser Phe Ser Arg Ala Ala Phe Asp Gly
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Pro Asn Arg Gln Ser Ala Arg Thr Leu Trp Tyr His Asp His Ala Met
165 170 175
His Ile Thr Ala Glu Asn Ala Tyr Arg Gly Gln Ala Gly Leu Tyr Met
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Leu Thr Asp Pro Ala Glu Asp Ala Leu Asn Leu Pro Ser Gly Tyr Gly
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Glu Phe Asp Ile Pro Met Ile Leu Thr Ser Lys Gln Tyr Thr Ala Asn
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Gly Asn Leu Val Thr Thr Asn Gly Glu Leu Asn Ser Phe Trp Gly Asp
225 230 235 240
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Arg Lys Tyr Arg Phe Arg Phe Leu Asp Ala Ala Val Ser Arg Ser Phe
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Gly Leu Tyr Phe Ala Asp Thr Asp Ala Ile Asp Thr Arg Leu Pro Phe
275 280 285
Lys Val Ile Ala Ser Asp Ser Gly Leu Leu Glu His Pro Ala Asp Thr
290 295 300
Ser Leu Leu Tyr Ile Ser Met Ala Glu Arg Tyr Glu Val Val Phe Asp
305 310 315 320
Phe Ser Asp Tyr Ala Gly Lys Thr Ile Glu Leu Arg Asn Leu Gly Gly
325 330 335
Ser Ile Gly Gly Ile Gly Thr Asp Thr Asp Tyr Asp Asn Thr Asp Lys
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Asn Arg Gln Phe Arg Phe Gly Arg Thr Gly Pro Thr Trp Thr Ile Asn
385 390 395 400
Gly Val Ala Phe Ala Asp Val Gln Asn Arg Leu Leu Ala Asn Val Pro
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Ala Pro Phe Pro Gly Val Tyr Met Phe His Cys His Asn Leu Ile His
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Glu Asp His Asp Met Met Ala Ala Phe Asn Ala Thr Val Leu Pro Asp
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Tyr Gly Tyr Asn Ala Thr Val Phe Val Asp Pro Met Glu Glu Leu Trp
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Gln Ala Arg Pro Tyr Glu Leu Gly Glu Phe Gln Ala Gln Ser Gly Gln
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